

1653



1600

p#16

RAW SEQUENCE LISTING

DATE: 10/17/2002

PATENT APPLICATION: US/09/043,944

TIME: 10:07:23

Input Set : A:\48231A_pctus_SEQLIST.ST25.txt

Output Set: N:\CRF4\10172002\I043944.raw

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OCT 25 2002

TECH CENTER 1600/2900

ENTERED

3 <110> APPLICANT: Greenwald, Iva
 4 Levitan, Diane
 6 <120> TITLE OF INVENTION: IDENTIFICATION OF SEL 12 AND USES THEREOF
 8 <130> FILE REFERENCE: 0575/48231-A-PCT-US
 10 <140> CURRENT APPLICATION NUMBER: 09/043,944
 11 <141> CURRENT FILING DATE: 1998-03-27
 13 <150> PRIOR APPLICATION NUMBER: PCT/US96/15727
 14 <151> PRIOR FILING DATE: 1996-09-27
 16 <150> PRIOR APPLICATION NUMBER: 60/004,387
 17 <151> PRIOR FILING DATE: 1995-09-27
 19 <160> NUMBER OF SEQ ID NOS: 27
 21 <170> SOFTWARE: PatentIn version 3.1
 23 <210> SEQ ID NO: 1
 24 <211> LENGTH: 461
 25 <212> TYPE: PRT
 26 <213> ORGANISM: C. elegans
 28 <220> FEATURE:
 29 <221> NAME/KEY: misc_feature
 30 <223> OTHER INFORMATION: Fig. 1A
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 36 1 5 10 15
 39 Thr His Thr Val Tyr Gly Thr Asn Leu Ile Thr Asn Arg Asn Ser Gln
 40 20 25 30
 43 Glu Asp Glu Asn Val Val Glu Ala Glu Leu Lys Tyr Gly Ala Ser
 44 35 40 45
 47 His Val Ile His Leu Phe Val Pro Val Ser Leu Cys Met Ala Leu Val
 48 50 55 60
 51 Val Phe Thr Met Asn Thr Ile Thr Phe Tyr Ser Gln Asn Asn Gly Arg
 52 65 70 75 80
 55 His Leu Leu Ser His Pro Phe Val Arg Glu Thr Asp Ser Ile Val Glu
 56 85 90 95
 59 Lys Gly Leu Met Ser Leu Gly Asn Ala Leu Val Met Leu Cys Val Val
 60 100 105 110
 63 Val Leu Met Thr Val Leu Leu Ile Val Phe Tyr Lys Tyr Lys Phe Tyr
 64 115 120 125
 67 Lys Leu Ile His Gly Trp Leu Ile Val Ser Ser Phe Leu Leu Leu Phe
 68 130 135 140
 71 Leu Phe Thr Thr Ile Tyr Val Gln Glu Val Leu Lys Ser Phe Asp Val
 72 145 150 155 160
 75 Ser Pro Ser Ala Leu Leu Val Leu Phe Gly Leu Gly Asn Tyr Gly Val
 76 165 170 175
 79 Leu Gly Met Met Cys Ile His Trp Lys Gly Pro Leu Arg Leu Gln Gln

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80          180          185          190
83 Phe Tyr Leu Ile Thr Met Ser Ala Leu Met Ala Leu Val Phe Ile Lys
84          195          200          205
87 Tyr Leu Pro Glu Trp Thr Val Trp Phe Val Leu Phe Val Ile Ser Val
88          210          215          220
91 Trp Asp Leu Val Ala Val Leu Thr Pro Lys Gly Pro Leu Arg Tyr Leu
92 225          230          235          240
95 Val Glu Thr Ala Gln Glu Arg Asn Glu Pro Ile Phe Pro Ala Leu Ile
96          245          250          255
99 Tyr Ser Ser Gly Val Ile Tyr Pro Tyr Val Leu Val Thr Ala Val Glu
100          260          265          270
103 Asn Thr Thr Asp Pro Arg Glu Pro Thr Ser Ser Asp Ser Asn Thr Ser
104          275          280          285
107 Thr Ala Phe Pro Gly Glu Ala Ser Cys Ser Ser Glu Thr Pro Lys Arg
108          290          295          300
111 Pro Lys Val Lys Arg Ile Pro Gln Lys Val Gln Ile Glu Ser Asn Thr
112 305          310          315          320
115 Thr Ala Ser Thr Thr Gln Asn Ser Gly Val Arg Val Glu Arg Glu Leu
116          325          330          335
119 Ala Ala Glu Arg Pro Thr Val Gln Asp Ala Asn Phe His Arg His Glu
120          340          345          350
123 Glu Glu Glu Arg Gly Val Lys Leu Gly Leu Gly Asp Phe Ile Phe Tyr
124          355          360          365
127 Ser Val Leu Leu Gly Lys Ala Ser Ser Tyr Phe Asp Trp Asn Thr Thr
128          370          375          380
131 Ile Ala Cys Tyr Val Ala Ile Leu Ile Gly Leu Cys Phe Thr Leu Val
132 385          390          395          400
135 Leu Leu Ala Val Phe Lys Arg Ala Leu Pro Ala Leu Gln Phe Pro Phe
136          405          410          415
139 Ser Pro Asp Ser Phe Phe Thr Phe Val Pro Ala Gly Ser Ser Pro His
140          420          425          430
143 Leu Leu His Lys Ser Leu Lys Ser Val Tyr Tyr Ile Asn Ser Leu Phe
144          435          440          445
147 Leu Pro Phe Leu Cys Ile Ile Asn Phe Ser Ile Ile Ser
148          450          455          460
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152 <211> LENGTH: 467
153 <212> TYPE: PRT
154 <213> ORGANISM: human
156 <220> FEATURE:
157 <221> NAME/KEY: misc_feature
158 <223> OTHER INFORMATION: Fig. 2A S182
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164 1          5          10          15
167 Ser Glu Asp Asn His Leu Ser Asn Thr Val Arg Ser Gln Asn Asp Asn
168          20          25          30
171 Arg Glu Arg Gln Glu His Asn Asp Arg Arg Ser Leu Gly His Pro Glu
172          35          40          45

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175 Pro Leu Ser Asn Gly Arg Pro Gln Gly Asn Ser Arg Gln Val Val Glu
176      50                      55                      60
179 Gln Asp Glu Glu Glu Asp Glu Glu Leu Thr Leu Lys Tyr Gly Ala Lys
180 65                      70                      75                      80
183 His Val Ile Met Leu Phe Val Pro Val Thr Leu Cys Met Val Val Val
184                      85                      90                      95
187 Val Ala Thr Ile Lys Ser Val Ser Phe Tyr Thr Arg Lys Asp Gly Gln
188                      100                      105                      110
191 Leu Ile Tyr Thr Pro Phe Thr Glu Asp Thr Glu Thr Val Gly Gln Arg
192                      115                      120                      125
195 Ala Leu His Ser Ile Leu Asn Ala Ala Ile Met Ile Ser Val Ile Val
196      130                      135                      140
199 Val Met Thr Ile Leu Leu Val Val Leu Tyr Lys Tyr Arg Cys Tyr Lys
200 145                      150                      155                      160
203 Val Ile His Ala Trp Leu Ile Ile Ser Ser Leu Leu Leu Phe Phe
204                      165                      170                      175
207 Phe Ser Phe Ile Tyr Leu Gly Glu Val Phe Lys Thr Tyr Asn Val Ala
208                      180                      185                      190
211 Val Asp Tyr Val Thr Val Ala Leu Leu Ile Trp Asn Phe Gly Val Val
212                      195                      200                      205
215 Gly Met Ile Ser Ile His Trp Lys Gly Pro Leu Arg Leu Gln Gln Ala
216      210                      215                      220
219 Tyr Leu Ile Met Ile Ser Ala Leu Met Ala Leu Val Phe Ile Lys Tyr
220 225                      230                      235                      240
223 Leu Pro Glu Trp Thr Ala Trp Leu Ile Leu Ala Val Ile Ser Val Tyr
224                      245                      250                      255
227 Asp Leu Val Ala Val Leu Cys Pro Lys Gly Pro Leu Arg Met Leu Val
228                      260                      265                      270
231 Glu Thr Ala Gln Glu Arg Asn Glu Thr Leu Phe Pro Ala Leu Ile Tyr
232                      275                      280                      285
235 Ser Ser Thr Met Val Trp Leu Val Asn Met Ala Glu Gly Asp Pro Glu
236      290                      295                      300
239 Ala Gln Arg Arg Val Ser Lys Asn Ser Lys Tyr Asn Ala Glu Ser Thr
240 305                      310                      315                      320
243 Glu Arg Glu Ser Gln Asp Thr Val Ala Glu Asn Asp Asp Gly Gly Phe
244                      325                      330                      335
247 Ser Glu Glu Trp Glu Ala Gln Arg Asp Ser His Leu Gly Pro His Arg
248                      340                      345                      350
251 Ser Thr Pro Glu Ser Arg Ala Ala Val Gln Glu Leu Ser Ser Ser Ile
252                      355                      360                      365
255 Leu Ala Gly Glu Asp Pro Glu Glu Arg Gly Val Lys Leu Gly Leu Gly
256      370                      375                      380
259 Asp Phe Ile Phe Tyr Ser Val Leu Val Gly Lys Ala Ser Ala Thr Ala
260 385                      390                      395                      400
263 Ser Gly Asp Trp Asn Thr Thr Ile Ala Cys Phe Val Ala Ile Leu Ile
264                      405                      410                      415
267 Gly Leu Cys Leu Thr Leu Leu Leu Leu Ala Ile Phe Lys Lys Ala Leu
268                      420                      425                      430
271 Pro Ala Leu Pro Ile Ser Ile Thr Phe Gly Leu Val Phe Tyr Phe Ala

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272          435          440          445
275 Thr Asp Tyr Leu Val Gln Pro Phe Met Asp Gln Leu Ala Phe His Gln
276          450          455          460
279 Phe Tyr Ile
280 465
283 <210> SEQ ID NO: 3
284 <211> LENGTH: 157
285 <212> TYPE: PRT
286 <213> ORGANISM: C. elegans
288 <220> FEATURE:
289 <221> NAME/KEY: MISC_FEATURE
290 <222> LOCATION: (11)..(11)
291 <223> OTHER INFORMATION: unknown amino acid
294 <400> SEQUENCE: 3
W--> 296 Glu Gly Lys Ser Pro Ser Asn Thr Glu Arg Xaa Val Ile Met Leu Phe
297 1          5          10          15
300 Val Pro Val Thr Leu Cys Met Ile Val Val Val Ala Thr Ile Lys Ser
301          20          25          30
304 Val Arg Phe Tyr Thr Glu Lys Asn Gly Gln Leu Ile Tyr Thr Pro Phe
305          35          40          45
308 Thr Glu Asp Thr Pro Ser Val Gly Gln Arg Leu Leu Asn Ser Val Leu
309          50          55          60
312 Asn Thr Leu Ile Met Ile Ser Val Ile Val Val Met Thr Ile Phe Leu
313 65          70          75          80
316 Val Val Leu Tyr Lys Tyr Arg Cys Tyr Lys Phe Ile His Gly Trp Leu
317          85          90          95
320 Ile Met Ser Ser Leu Met Leu Leu Phe Leu Phe Thr Tyr Ile Tyr Leu
321          100          105          110
324 Gly Glu Val Leu Lys Thr Tyr Asn Val Ala Met Asp Tyr Pro Thr Leu
325          115          120          125
328 Leu Leu Thr Val Trp Glu Leu Arg Gly Ser Gly His Gly Val His Pro
329          130          135          140
332 Leu Glu Gly Ala Phe Gly Ala Ala Glu Ala Tyr Leu Ser
333 145          150          155
336 <210> SEQ ID NO: 4
337 <211> LENGTH: 465
338 <212> TYPE: PRT
339 <213> ORGANISM: C. elegans
341 <220> FEATURE:
342 <221> NAME/KEY: misc_feature
343 <223> OTHER INFORMATION: Fig. 2A SPE-4
346 <400> SEQUENCE: 4
348 Met Asp Thr Leu Arg Ser Ile Ser Ser Glu Leu Val Arg Ser Ser Gln
349 1          5          10          15
352 Leu Arg Trp Thr Leu Phe Ser Val Ile Ala Asn Met Ser Leu Thr Leu
353          20          25          30
356 Ser Ile Trp Ile Gly Val Tyr Asn Met Glu Val Asn Ser Glu Leu Ser
357          35          40          45
360 Lys Thr Tyr Phe Leu Asp Pro Ser Phe Glu Gln Thr Thr Gly Asn Leu

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Input Set : A:\48231A_pctus_SEQLIST.ST25.txt

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361      50      55      60
364 Leu Leu Asp Gly Phe Ile Asn Gly Val Gly Thr Ile Leu Val Leu Gly
365 65      70      75      80
368 Cys Val Ser Phe Ile Met Leu Ala Phe Val Leu Phe Asp Phe Arg Arg
369      85      90      95
372 Ile Val Lys Ala Trp Leu Thr Leu Ser Cys Leu Leu Ile Leu Phe Gly
373      100      105      110
376 Val Ser Ala Gln Thr Leu His Asp Met Phe Ser Gln Val Phe Asp Gln
377      115      120      125
380 Asp Asp Asn Asn Gln Tyr Tyr Met Thr Ile Val Leu Ile Val Val Pro
381      130      135      140
384 Thr Val Val Tyr Gly Phe Gly Gly Ile Tyr Ala Phe Phe Ser Asn Ser
385 145      150      155      160
388 Ser Leu Ile Leu His Gln Ile Phe Val Val Thr Asn Cys Ser Leu Ile
389      165      170      175
392 Ser Val Phe Tyr Leu Arg Val Phe Pro Ser Lys Thr Thr Trp Phe Val
393      180      185      190
396 Leu Trp Ile Val Leu Phe Trp Asp Leu Phe Ala Val Leu Ala Pro Met
397      195      200      205
400 Gly Pro Leu Lys Lys Val Gln Glu Lys Ala Ser Asp Tyr Ser Lys Cys
401      210      215      220
404 Val Leu Asn Leu Ile Met Phe Ser Ala Asn Glu Lys Arg Leu Thr Ala
405 225      230      235      240
408 Gly Ser Asn Gln Glu Glu Thr Asn Glu Gly Glu Glu Ser Thr Ile Arg
409      245      250      255
412 Arg Thr Val Lys Gln Thr Ile Glu Tyr Tyr Thr Lys Arg Glu Ala Gln
413      260      265      270
416 Asp Asp Glu Phe Tyr Gln Lys Ile Arg Gln Arg Arg Ala Ala Ile Asn
417      275      280      285
420 Pro Asp Ser Val Pro Thr Glu His Ser Pro Leu Val Glu Ala Glu Pro
421      290      295      300
424 Ser Pro Ile Glu Leu Lys Glu Lys Asn Ser Thr Glu Glu Leu Ser Asp
425 305      310      315      320
428 Asp Glu Ser Asp Thr Ser Glu Thr Ser Ser Gly Ser Ser Asn Leu Ser
429      325      330      335
432 Ser Ser Asp Ser Ser Thr Thr Val Ser Thr Ser Asp Ile Ser Thr Ala
433      340      345      350
436 Glu Glu Cys Asp Gln Lys Glu Trp Asp Asp Leu Val Ser Asn Ser Leu
437      355      360      365
440 Pro Asn Asn Asp Lys Arg Pro Ala Thr Ala Ala Asp Ala Leu Asn Asp
441      370      375      380
444 Gly Glu Val Leu Arg Leu Gly Phe Gly Asp Phe Val Phe Tyr Ser Leu
445 385      390      395      400
448 Leu Ile Gly Gln Ala Ala Ala Ser Gly Cys Pro Phe Ala Val Ile Ser
449      405      410      415
452 Ala Ala Leu Gly Ile Leu Phe Gly Leu Val Val Thr Leu Thr Val Phe
453      420      425      430
456 Ser Thr Glu Glu Ser Thr Thr Pro Ala Leu Pro Leu Pro Val Ile Cys
457      435      440      445

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 10/17/2002
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Input Set : A:\48231A_pctus_SEQLIST.ST25.txt
Output Set: N:\CRF4\10172002\I043944.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 11